

Serial Number: 09/699,652A

Errors Corrected by the STIC Systems Branch

CRF Processing Date: 11/4/2002  
Edited by: AE  
Verified by: (STIC staff)

1600

11/4/2002

#14

Changed a file from non-ASCII to ASCII

**ENTERED**

Changed the margins in cases where the sequence text was wrapped down to the next line.

Edited a format error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was  the prior application data; or  other \_\_\_\_\_.

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included:

**RECEIVED**

NOV 07 2002

Deleted extra, invalid, headings used by an applicant, specifically:

**TECH CENTER 1600/2900**

Deleted:  non-ASCII "garbage" at the beginning/end of files;  secretary initials/filename at end of file;  
 page numbers throughout text;  other invalid text, such as \_\_\_\_\_.

Inserted mandatory headings, specifically:

Corrected an obvious error in the response, specifically:

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_

Other:

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



1600

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/699,652A**

**DATE: 11/04/2002**  
**TIME: 17:50:29**

**Input Set : A:\PTO.AMC.txt**  
**Output Set: N:\CRF4\11042002\I699652A.raw**

P.6

3 <110> APPLICANT: Cahoon, Edgar B.  
 4       Cahoon, Rebecca E.  
 5       Kinney, Anthony J.  
 6       Rafalski, J. Antoni  
 8 <120> TITLE OF INVENTION: TRIACYLGLYCEROL LIPASES  
 10 <130> FILE REFERENCE: BB1168 US NA  
 12 <140> CURRENT APPLICATION NUMBER: 09/699,652A  
 13 <141> CURRENT FILING DATE: 2002-10-30  
 15 <150> PRIOR APPLICATION NUMBER: 60/083,688  
 16 <151> PRIOR FILING DATE: 1988-04-30  
 18 <150> PRIOR APPLICATION NUMBER: PCT/US99/09280  
 19 <151> PRIOR FILING DATE: 1999-04-29  
 21 <160> NUMBER OF SEQ ID NOS: 36  
 23 <170> SOFTWARE: Microsoft Office 97  
 25 <210> SEQ ID NO: 1  
 26 <211> LENGTH: 751  
 27 <212> TYPE: DNA  
 28 <213> ORGANISM: Zea mays  
 30 <400> SEQUENCE: 1  
 31 gcacgagatc accggcaaga actactgcct caacagctcc gccgtcgacg tcttcctcaa 60  
 32 gtacgagccc cagccgaccc ccaccaaaac catggtccac ttgcgtcaaa ccgtgcgcga 120  
 33 cggcgtgctg accaagtacg actacgtgct gccggagcgg aacatcgcca gctacggcca 180  
 34 ggccgagccg ccgggtgtacc ggtatgtccgg catcccgccg agcttccgcg tcttcctcag 240  
 35 ctacggcggc cgggactcgc tcgcccggacc cggccgacgtg cgcctccccc tgcaaggacct 300  
 36 ccggggccac gaccaggaca agtcacgggt gcagttacccg gacaaggatcg cgcacccctoga 360  
 37 cttcatcatc ggcgtctgct ccaaggacta cgtctacaag gacatgtatcg actttctaaa 420  
 38 ccgttcaac tagtacttagc atatatattt gttcaatcg gtgtcgctt cagccccagc 480  
 39 aggatttagac aaaaaaaaggg ggggacactg cagctcgtaa acgttgtcca tacagattat 540  
 40 cagaggtgaa aaccatacat gatgttaattt agcatttagat agttaaaaca tggagctgcc 600  
 41 tcagtatggg ggattgtcaa ctactctcca tcacagcagt aggtgtatcg tagaagagtg 660  
 42 attgtcacac tgtgtgtgtt gcaaatttgc aacacagtga ttactaatat aaaaaatact 720  
 43 cttgagttaa aaaaaaaaaa aaaaaaaaaa a 751  
 45 <210> SEQ ID NO: 2  
 46 <211> LENGTH: 143  
 47 <212> TYPE: PRT  
 48 <213> ORGANISM: Zea mays  
 50 <400> SEQUENCE: 2  
 51 His Glu Ile Thr Gly Lys Asn Tyr Cys Leu Asn Ser Ser Ala Val Asp  
 52     1                   5                   10                   15  
 54 Val Phe Leu Lys Tyr Glu Pro Gln Pro Thr Ser Thr Lys Thr Met Val  
 55     20                25                30  
 57 His Phe Ala Gln Thr Val Arg Asp Gly Val Leu Thr Lys Tyr Asp Tyr  
 58     35                40                45

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Input Set : A:\PTO.AMC.txt  
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60 Val Leu Pro Glu Arg Asn Ile Ala Ser Tyr Gly Gln Ala Glu Pro Pro  
61 50 55 60  
63 Val Tyr Arg Met Ser Gly Ile Pro Pro Ser Phe Pro Leu Phe Leu Ser  
64 65 70 75 80  
66 Tyr Gly Gly Arg Asp Ser Leu Ala Asp Pro Ala Asp Val Arg Leu Leu  
67 85 90 95  
69 Leu Gln Asp Leu Arg Gly His Asp Gln Asp Lys Leu Thr Val Gln Tyr  
70 100 105 110  
72 Leu Asp Lys Phe Ala His Leu Asp Phe Ile Ile Gly Val Cys Ala Lys  
73 115 120 125  
75 Asp Tyr Val Tyr Lys Asp Met Ile Asp Phe Leu Asn Arg Phe Asn  
76 130 135 140  
78 <210> SEQ ID NO: 3  
79 <211> LENGTH: 647  
80 <212> TYPE: DNA  
81 <213> ORGANISM: Catalpa sp.  
83 <400> SEQUENCE: 3  
84 ttatcttca ggagagattt ttgtttgaat gtcggcccg ttgagcttt tggaaaaat 60  
85 taccctccat cttccgtgaa ttgagacccc tgtccatatg gctcaaactg tcgcataatgg 120  
86 gatcctaccc aaatacact acggcaatcc cagctcaac ttggcccatt atggtaatc 180  
87 cagacctccc gtttacatt tatccaagat tccccctcgac attccgctct tcctaagcta 240  
88 tggaggacaa gatgcattgt cgatgttaa ggatgtcgag acattgctcg atagtctcaa 300  
89 gttacacgt gtggataaagc tgcgtgtca gtataatcaag gattatgctc atgccgactt 360  
90 cattatcgga gttactgcaa aagatatagt ttataatcag attgttaactt ttttcagaaa 420  
91 ccaggcttga gaggttcttg attttggagt gctttgctg tgagaatgca acagttgtt 480  
92 ccactcttgt tgaatgtgaa taagccattt ccgagagatt taatggctgg taaagcttat 540  
93 tagttactc atagatacat gtaagaagca acccgataca tagttgaat ccttatctc 600  
94 gaaaaggtat tgcacatccct ctctacgtc aaaaaaaaaaaa aaaaata 647  
96 <210> SEQ ID NO: 4  
97 <211> LENGTH: 116  
98 <212> TYPE: PRT  
99 <213> ORGANISM: Catalpa sp.  
101 <400> SEQUENCE: 4  
102 Ile Glu Thr Pro Val His Met Ala Gln Thr Val Arg Tyr Gly Ile Leu  
103 1 5 10 15  
105 Pro Lys Tyr Asp Tyr Gly Asn Pro Ser Phe Asn Leu Ala His Tyr Gly  
106 20 25 30  
108 Glu Ser Arg Pro Pro Val Tyr Asp Leu Ser Lys Ile Pro Leu Asp Ile  
109 35 40 45  
111 Pro Leu Phe Leu Ser Tyr Gly Gly Gln Asp Ala Leu Ser Asp Val Lys  
112 50 55 60  
114 Asp Val Glu Thr Leu Leu Asp Ser Leu Lys Leu His Asp Val Asp Lys  
115 65 70 75 80  
117 Leu His Val Gln Tyr Ile Lys Asp Tyr Ala His Ala Asp Phe Ile Ile  
118 85 90 95  
120 Gly Val Thr Ala Lys Asp Ile Val Tyr Asn Gln Ile Val Thr Phe Phe  
121 100 105 110  
123 Arg Asn Gln Ala  
124 115

RAW SEQUENCE LISTING  
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Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\11042002\I699652A.raw

126 <210> SEQ ID NO: 5  
 127 <211> LENGTH: 705  
 128 <212> TYPE: DNA  
 129 <213> ORGANISM: Catalpa sp.  
 131 <220> FEATURE:  
 132 <221> NAME/KEY: unsure  
 133 <222> LOCATION: (526) /  
 134 <223> OTHER INFORMATION: n = A, C, G, or T  
 136 <220> FEATURE:  
 137 <221> NAME/KEY: unsure  
 138 <222> LOCATION: (561) /  
 139 <223> OTHER INFORMATION: n = A, C, G, or T  
 141 <220> FEATURE:  
 142 <221> NAME/KEY: unsure  
 143 <222> LOCATION: (585) /  
 144 <223> OTHER INFORMATION: n = A, C, G, or T  
 146 <220> FEATURE:  
 147 <221> NAME/KEY: unsure  
 148 <222> LOCATION: (593) /  
 149 <223> OTHER INFORMATION: n = A, C, G, or T  
 151 <220> FEATURE:  
 152 <221> NAME/KEY: unsure  
 153 <222> LOCATION: (664) /  
 154 <223> OTHER INFORMATION: n = A, C, G, or T  
 156 <220> FEATURE:  
 157 <221> NAME/KEY: unsure  
 158 <222> LOCATION: (679) /  
 159 <223> OTHER INFORMATION: n = A, C, G, or T  
 161 <400> SEQUENCE: 5  
 162 gcacgagcca acagcttccat aattttagct ttcttaatcc ttctctcatt atcactactc 60  
 163 ctacctcattc aatcatttcgc ctccagccgc cggcgttttc ttccgcagaa tggatgtcggtt 120  
 164 ctccgcggc acggcggttg ctccaccggc gtaactgtac acggttataa atgccaagaa 180  
 165 tttgaagtttca cgactgtatgtatata ttaagcgtgc agaggatct ggaggggccgg 240  
 166 gccggaggag gagggccgaa gcggccgcgc gttctgtgc aacatgggggt tcttgtggac 300  
 167 gggatgacgt ggtctgtgaa tggaccggaa caatcttgg cgatgtatatt ggctgacaat 360  
 168 gggttcgacg tctggatttc taacataaga ggaacttaggt ttagtcgtcg tcatgtcagc 420  
 169 ctgtatcccta ccgatctga atattggat tggcatggg acgtatctgg tgacccacga 480  
 W--> 170 cttagccatcc ctgtatcgagt tagtggtcag acaaacgggt cagaanacac actacatagg 540  
 W--> 171 gcaatccatg gggaaacctta nttagctttgg gatcaactttt aggangggaaa cangttggca 600  
 172 gggtaaatcg gctgtatgtt aagccaaatgg gctaacgagt catatgcaac tgctctcgag 660  
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 175 <210> SEQ ID NO: 6  
 176 <211> LENGTH: 157  
 177 <212> TYPE: PRT  
 178 <213> ORGANISM: Catalpa sp.  
 180 <400> SEQUENCE: 6  
 181 Ala Arg Ala Asn Ser Phe Leu Asn Leu Ala Leu Ile Leu Leu Ser  
 182 1 5 10 15  
 184 Leu Ser Leu Leu Pro His Gln Ser Phe Ala Ser Ser Arg Arg Arg

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/699,652A

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Input Set : A:\PTO.AMC.txt  
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```

185          20          25          30
187 Phe Leu Pro Gln Asn Asp Val Val Leu Pro Pro Asp Gly Val Cys Ser
188          35          40          45
190 Thr Ala Val Thr Val His Gly Tyr Lys Cys Gln Glu Phe Glu Val Thr
191          50          55          60
193 Thr Asp Asp Gly Tyr Ile Leu Ser Val Gln Arg Ile Leu Glu Gly Arg
194          65          70          75          80
196 Ala Gly Gly Gly Pro Lys Arg Pro Pro Val Leu Leu Gln His Gly
197          85          90          95
199 Val Leu Val Asp Gly Met Thr Trp Leu Val Asn Gly Pro Glu Gln Ser
200          100         105         110
202 Leu Ala Met Ile Leu Ala Asp Asn Gly Phe Asp Val Trp Ile Ser Asn
203          115         120         125
205 Ile Arg Gly Thr Arg Phe Ser Arg Arg His Val Ser Leu Asp Pro Thr
206          130         135         140
208 Asp Pro Glu Tyr Trp Asp Trp Ala Trp Asp Asp Leu Gly
209 145          150          155
211 <210> SEQ ID NO: 7
212 <211> LENGTH: 859
213 <212> TYPE: DNA
214 <213> ORGANISM: Zea mays
216 <220> FEATURE:
217 <221> NAME/KEY: unsure
218 <222> LOCATION: (46)
219 <223> OTHER INFORMATION: n = A, C, G, or T
221 <220> FEATURE:
222 <221> NAME/KEY: unsure
223 <222> LOCATION: (231)
224 <223> OTHER INFORMATION: n = A, C, G, or T
226 <400> SEQUENCE: 7
W--> 227 aaagcaaaca acggccgaca tggtgccccc aggaaaagcg cttgcngcgc cccagctcct 60
228 ctcctcggt ttccctgcc tcctagccgg tggagccgc gcatccccgc ccacagacgc 120
229 gtcacgcccgc gtctccccgc gcgcgggggc cgggtggctc tgccagcagc tgccctgcc 180
W--> 230 gcagggttacc cgtgcaccga gcacaccgtt caaacggatg atggctttct nttgtctctt 240
231 cagcatattc cacatggcag aatggaaatt gcagataata ctggacctcc agttttctt 300
232 cagcacggtc tttccaggg tggagataca tggttcataa actccaatga acaatcaactt 360
233 ggatatatcc ttgctgacaa tggttttgtat gtttgggtcg gaaatgtcg tggcacacgt 420
234 tggagtaaag gccactctac tctctctgtt catgataagc ttttctggta ttggagttgg 480
235 caagaccttg ctgaatacga cttttggca atgttaagct atgtatatac agttgcacag 540
236 tccaaaattt tgtatgtggg acattcacag ggaactatca tgggttggc tgcgtttaca 600
237 atgcctgaaa cagtaaagat gataagctt gctgcgttc tttgtcccat ttcttacctt 660
238 gatcacgtca gtgctagtt tggctttaga gcagttgcca tgcatttcttga tgagatgttt 720
239 gttattatgg gcatccatca gttgaacttc cggagcgata tgggtgtaca gatatttagat 780
240 tcgctgtgcg atgatgaaca tttggactgc aacgatctgt tatcttcaat aacagtcaaa 840
241 actgttgttc aatcatctc                                     859
243 <210> SEQ ID NO: 8
244 <211> LENGTH: 286
245 <212> TYPE: PRT
246 <213> ORGANISM: Zea mays

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RAW SEQUENCE LISTING  
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Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\11042002\I699652A.raw

248 <220> FEATURE:  
 249 <221> NAME/KEY: UNSURE  
 250 <222> LOCATION: (16)  
 251 <223> OTHER INFORMATION: ANY AMINO ACID  
 253 <400> SEQUENCE: 8  
 W--> 254 Lys Ala Asn Asn Gly Gly His Gly Ala Pro Arg Lys Ser Ala Cys Xaa  
 255 1 5 10 15  
 257 Ala Pro Ala Pro Pro Pro Arg Val Pro Leu Pro Pro Ser Arg Trp Ser  
 258 20 25 30  
 260 Pro Arg Ile Pro Ala His Arg Arg Ala Thr Pro Arg Leu Pro Ala Arg  
 261 35 40 45  
 263 Gly Gly Arg Trp Pro Leu Pro Ala Ala Ala Pro Ala Ala Gly Tyr Pro  
 264 50 55 60  
 266 Cys Thr Glu His Thr Val Gln Thr Asp Asp Gly Phe Leu Leu Ser Leu  
 267 65 70 75 80  
 269 Gln His Ile Pro His Gly Arg Asn Gly Ile Ala Asp Asn Thr Gly Pro  
 270 85 90 95  
 272 Pro Val Phe Leu Gln His Gly Leu Phe Gln Gly Gly Asp Thr Trp Phe  
 273 100 105 110  
 275 Ile Asn Ser Asn Glu Gln Ser Leu Gly Tyr Ile Leu Ala Asp Asn Gly  
 276 115 120 125  
 278 Phe Asp Val Trp Val Gly Asn Val Arg Gly Thr Arg Trp Ser Lys Gly  
 279 130 135 140  
 281 His Ser Thr Leu Ser Val His Asp Lys Leu Phe Trp Asp Trp Ser Trp  
 282 145 150 155 160  
 284 Gln Asp Leu Ala Glu Tyr Asp Val Leu Ala Met Leu Ser Tyr Val Tyr  
 285 165 170 175  
 287 Thr Val Ala Gln Ser Lys Ile Leu Tyr Val Gly His Ser Gln Gly Thr  
 288 180 185 190  
 290 Ile Met Gly Leu Ala Ala Phe Thr Met Pro Glu Thr Val Lys Met Ile  
 291 195 200 205  
 293 Ser Ser Ala Ala Leu Leu Cys Pro Ile Ser Tyr Leu Asp His Val Ser  
 294 210 215 220  
 296 Ala Ser Phe Val Leu Arg Ala Val Ala Met His Leu Asp Glu Met Leu  
 297 225 230 235 240  
 299 Val Ile Met Gly Ile His Gln Leu Asn Phe Arg Ser Asp Met Gly Val  
 300 245 250 255  
 302 Gln Ile Leu Asp Ser Leu Cys Asp Asp Glu His Leu Asp Cys Asn Asp  
 303 260 265 270  
 305 Leu Leu Ser Ser Ile Thr Val Lys Thr Val Val Gln Ser Ser  
 306 275 280 285  
 308 <210> SEQ ID NO: 9  
 309 <211> LENGTH: 509  
 310 <212> TYPE: DNA  
 311 <213> ORGANISM: Zea mays  
 313 <220> FEATURE:  
 314 <221> NAME/KEY: unsure  
 315 <222> LOCATION: (162)  
 316 <223> OTHER INFORMATION: n = A, C, G, or T

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/699,652A

DATE: 11/04/2002  
TIME: 17:50:30

Input Set : A:\PTO.AMC.txt  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 526,561,585,593,664,679  
Seq#:7; N Pos. 46,231  
Seq#:8; Xaa Pos. 16  
Seq#:9; N Pos. 162,277,284,290,295,386,406,413,443,468,484,489  
Seq#:10; Xaa Pos. 52,90,92,96  
Seq#:11; N Pos. 8,20,229,236,241,249,268  
Seq#:12; Xaa Pos. 76,78,80,83,89  
Seq#:15; N Pos. 12,24,29,33,43,78,182,265,300,302,306,347,351,367,370,380  
Seq#:15; N Pos. 386  
Seq#:16; Xaa Pos. 8,10,11,15,61  
Seq#:23; N Pos. 226,315,462,1306,1349,1359,1368,1373  
Seq#:24; Xaa Pos. 50,80,129  
Seq#:25; N Pos. 601  
Seq#:26; Xaa Pos. 45  
Seq#:27; N Pos. 7,15,27,38,50,94,99,103,105,117

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/699,652A

DATE: 11/04/2002  
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Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\11042002\I699652A.raw

L:170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:480  
L:171 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:540  
L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:660  
L:227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0  
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:180  
L:254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0  
L:376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:120  
L:378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:240  
L:380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:360  
L:381 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:420  
L:382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:480  
L:419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:48  
L:425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:80  
L:475 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0  
L:478 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:180  
L:479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:240  
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:64  
L:527 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:80  
L:737 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0  
L:738 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:60  
L:740 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:180  
L:741 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:240  
L:742 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:300  
L:743 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:360  
L:771 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0  
L:780 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:48  
L:1124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:180  
L:1126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:300  
L:1128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:420  
L:1142 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:1260  
L:1143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:1320  
L:1176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:48  
L:1179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:64  
L:1191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:128  
L:1245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:600  
L:1279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:32  
L:1380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0  
L:1381 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:60